

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/664,859A
Source: IFW/6
Date Processed by STIC: 5-26-06

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 05/26/2006

PATENT APPLICATION: US/10/664,859A

TIME: 12:16:26

Input Set : A:\Q77377 seq.ST25.txt

Output Set: N:\CRF4\05262006\J664859A.raw

3 <110> APPLICANT: BASLER, Konrad
 4 BRUNNER, Erich
 5 FROESCH, Barbara
 6 KRAMPS, Thomas
 7 PETER, Oliver
 9 <120> TITLE OF INVENTION: ESSENTIAL DOWNSTREAM COMPONENT OF THE WINGLESS SIGNALING
 PATHWAY
 10 AND THERAPEUTIC AND DIAGNOSTIC APPLICATIONS BASED THEREON
 12 <130> FILE REFERENCE: Q77377
 14 <140> CURRENT APPLICATION NUMBER: 10/664,859A
 15 <141> CURRENT FILING DATE: 2003-09-22
 17 <150> PRIOR APPLICATION NUMBER: US 09/915,543
 18 <151> PRIOR FILING DATE: 2001-07-27
 20 <150> PRIOR APPLICATION NUMBER: 60/221,502
 21 <151> PRIOR FILING DATE: 2000-07-28
 23 <160> NUMBER OF SEQ ID NOS: 25
 25 <170> SOFTWARE: PatentIn version 3.3
 27 <210> SEQ ID NO: 1
 28 <211> LENGTH: 6909
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 55 <222> LOCATION: (4927)..(6456)
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 62 ttagcttaat acgatgctcc gaagtgttat tgcatttgca catatacata aaattgtgac 180
 64 atagaatagg agaattccac atacaaatac aaaaatacaa aatcctccag taaaatttaa 240

(pg.6)

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70 agcaataaaa cgatggattt taattgctac ttgagcaatt agccacacaa gggatcttgg      420
72 gaaggtcgat ttgaaggaat tcgatttcta ggatgctctc gacaaca atg ccc cgc      476
73                                     Met Pro Arg
74                                     1
76 agt cca acc caa caa cag ccg caa cca aac tcc gat gcc tcc tca aca      524
77 Ser Pro Thr Gln Gln Gln Pro Gln Pro Asn Ser Asp Ala Ser Ser Thr
78      5                      10                      15
80 agt gca tct gga tca aat cct gga gca gcg atc gga aat ggg gac tcg      572
81 Ser Ala Ser Gly Ser Asn Pro Gly Ala Ala Ile Gly Asn Gly Asp Ser
82 20                      25                      30                      35
84 gcg gcg agc aga agt tct ccg aag acc ctt aat agc gaa ccc ttt tct      620
85 Ala Ala Ser Arg Ser Ser Pro Lys Thr Leu Asn Ser Glu Pro Phe Ser
86      40                      45                      50
88 act ttg tcg ccg ggtaagactt gtattgattt ctctttgtcc ggaattataa      672
89 Thr Leu Ser Pro
90      55
92 caactttctg tgtttcca gat caa ata aaa ttg acg cca gaa gaa ggc act      723
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96 gag aaa agc gga cta tca act agt gat aaa gct gcc act gga gga gcc      771
97 Glu Lys Ser Gly Leu Ser Thr Ser Asp Lys Ala Ala Thr Gly Gly Ala
98      70                      75                      80
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104 aac tct acg agc aca atc aac tcg tgc cta gtc gct tct cca caa aac      867
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110 115                      120                      125                      130
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116 gtg aag gac gag gaa gct ggtaagactg ccctacaaat ggtttaaaat      1011
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142 gtc aag gaa gaa ccc acc gat gtc tta ggc agt tta gta aat atg aaa      1578
143 Val Lys Glu Glu Pro Thr Asp Val Leu Gly Ser Leu Val Asn Met Lys
144          180          185          190
146 aaa gaa gaa aga gaa aat cat tcg cca acg atg tcc cct gtt ggt ttt      1626
147 Lys Glu Glu Arg Glu Asn His Ser Pro Thr Met Ser Pro Val Gly Phe
148          195          200          205
150 ggt tca att ggt aat gca cag gac aac tcc gct aca ccg ggtaagtttt      1675
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152 210          215          220
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156 tcaaagaact actatagcga tatctcctgc cttttaattt tattttaatt aggaaatacg      1795
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164 ggaaaaatggg tgtaattttg taggaagttt tcattttaga agaaatgtga ttattttatt      2035
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176 agttatttaa ttttttatgg tgtattttaa taaatatcct atttattcat ttacata      2393
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186 cag ttg aat ccc gat ttt atc aat gaa tct tta aat aat cct gca att      2537
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188 255          260          265          270
190 tcg agc ata tta gta agc gga gta gga cca ata ccc gga atc gga gtt      2585
191 Ser Ser Ile Leu Val Ser Gly Val Gly Pro Ile Pro Gly Ile Gly Val
192          275          280          285
194 gga gcg ggg acg gga aat tta ttg act gcc aac gcc aat gga atc tcc      2633
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196          290          295          300
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216					370					375						380	
218	ggg	cag	gtt	gga	cta	act	cct	cct	aat	cct	gta	gcc	aaa	ata	aca	caa	2921
219	Gly	Gln	Val	Gly	Leu	Thr	Pro	Pro	Asn	Pro	Val	Ala	Lys	Ile	Thr	Gln	
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222	cag	cag	cca	cat	aca	aag	acc	gta	ggc	cta	ttg	aaa	ccc	caa	ttc	aat	2969
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291	Ser	Phe	Gly	Asn	Thr	Ser	Cys	Asn	Val	Ile	Gly	Thr	Ala	Pro	Asp	Met													
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314	aga	tct	gcg	tca	gta	cca	ata	gcc	act	caa	tcg	ccc	aat	ccc	tcg	agt	4073												
315	Arg	Ser	Ala	Ser	Val	Pro	Ile	Ala	Thr	Gln	Ser	Pro	Asn	Pro	Ser	Ser													
316	770																775				780								
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328	815																820				825				830				
330	aca	gtg	ctc	tca	gca	aac	aag	aac	tgt	ttt	cag	gca	gac	acc	cca	tcg	4265												
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:24; Xaa Pos. ~~1,2,7,8,12,15,18,21,22,25,26,27~~
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Seq#:25; Xaa Pos. 31,32,33

VERIFICATION SUMMARY

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L:2203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0

M:341 Repeated in SeqNo=24

L:2249 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0

M:341 Repeated in SeqNo=25